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PubMed Nucleotide Protein Genome Structure PopSet

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1 : O65607 . DNA MISMATCH REPAI...[gi:3914062] Related Sequences

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LOCUS       MSH3_ARATH   1076 aa                    PLN       15-DEC-1998
DEFINITION  DNA MISMATCH REPAIR PROTEIN MSH3.
ACCESSION   O65607
PID         g3914062
VERSION     O65607  GI:3914062
DBSOURCE    swissprot: locus MSH3_ARATH, accession O65607;
            class: standard.
            created: Dec 15, 1998.
            sequence updated: Dec 15, 1998.
            annotation updated: Dec 15, 1998.
            xrefs: gi: 5596409, gi: 2980796
            xrefs (non-sequence databases): PFAM PF00488
KEYWORDS    DNA repair; ATP-binding; DNA-binding.
SOURCE      thale cress.
  ORGANISM  Arabidopsis thaliana
            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophy
            euphyllophytes; Spermatophyta; Magnoliophyta; eudicotyledons; c
            eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae;
            Arabidopsis.
REFERENCE   1  (residues 1 to 1076)
  AUTHORS   BEVAN,M., BARGUES,M., PEREZ-PEREZ,A., TEROL,J., TORRES,A.,
            PEREZ-ALONSO,M., HOHEISEL,J., MEWES,H.-W., MAYER,K. and
            SCHUELLER,C.
  TITLE     Direct Submission
  JOURNAL   Submitted (??-MAR-1998) to the EMBL/GenBank/DDBJ databases
  REMARK    SEQUENCE FROM N.A.
            STRAIN=CV. COLUMBIA
COMMENT     ---------------------------------------------------------------
            This SWISS-PROT entry is copyright. It is produced through a
            collaboration between the Swiss Institute of Bioinformatics and
            the EMBL outstation - the European Bioinformatics Institute.
            The original entry is available from http://www.expasy.ch/sprot
            and http://www.ebi.ac.uk/sprot
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            [FUNCTION] NOT KNOWN. PROBABLE DNA-REPAIR PROTEIN.
            [SIMILARITY] BELONGS TO DNA MISMATCH REPAIR MUTS FAMILY.
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Revised: January 10, 2000.

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ID   ATM7J2     standard; DNA; PLN; 80386 BP.
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AC   AL022197;
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SV   AL022197.2
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DT   18-MAR-1998 (Rel. 55, Created)
DT   23-JUL-1999 (Rel. 60, Last updated, Version 5)
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DE   Arabidopsis thaliana DNA chromosome 4, P1 clone M7J2 (ESSA project)
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OC   euphyllophytes; Spermatophyta; Magnoliophyta; eudicotyledons;
OC   core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae;
OC   Arabidopsis.
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RN   [1]
RA   Bevan M., Bargues M., Perez-Perez A., Terol J., Torres A., Perez-Alonso M.,
RA   Hoheisel J., Mewes H.W., Mayer K.F.X., Lemcke K., Schueller C.;
RT   ;
RL   Unpublished.
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RN   [2]
RP   1-80386
RA   EU Arabidopsis sequencing project;
RT   ;
RL   Submitted (23-JUL-1999) to the EMBL/GenBank/DDBJ databases.
RL   MIPS, at the Max-Planck-Institut fuer Biochemie, Am Klopferspitz 18a,
RL   D-82152 Martinsried, FRG, E-mail:
RL   schuelle@mips.biochem.mpg.de,mayer@mips.biochem.mpg.de Project Coordinator:
RL   Mike Bevan, Molecular Genetics Department, Cambridge Laboratory, John Innes
RL   Centre, Colney Lane, NR4 7UJ Norwich, UK, E-mail: michael.bevan@bbsrc.ac.uk
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DR   SWISS-PROT; O65607; MSH3_ARATH.
DR   SWISS-PROT; P92965; RS40_ARATH.
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CC   Information on performance of analysis and a more detailed annotation
CC   of this entry and other sequences of chromosomes 3, 4 and 5 can be
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CC   viewed at: http://www.mips.biochem.mpg.de/proj/thal/
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aagacaaaag acaaaaaaaa aaaaactaat caatatgtta attagattta taatataaaa  79860
```

```
     tttaggaatg attccaccac tgtttttgtc aacatcattc caccacttgt tttagcgaat     79920
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     acaacaaaaa aagaagtgga caacactata cgatcgactt aaatgcttgc ttatataaag     80100
     actaaaagga ccattggttc ccgtatctcc tcgcaatact actactcact ttactataat     80160
     ctctcaaaat ggccgtaagt ttcgtaacaa catctcctga ggaagaagac aaaccgaagc     80220
     taggccttgg aaatattcaa actccgttaa tcttcaaccc ttcaatgctt aaccttcaag     80280
     ccaatatccc aaaccaattc atctggcctg acgacgaaaa accttccatc aacgttctcg     80340
     agcttgatgt tcctctcatc gaccttcaaa accttctctc tgatcc                    80386
//
```